

Table S1.**Primers used in the expression and functional analysis of the *SLC27A1* promoter**

Reaction	primer	Primer sequence(5'-3')	Binding region	Size (bp)
qRT-PCR	FATP1 forward	AAGAGCCTGGTCAAGTTCTG	--	239
	FATP1 reverse	TAGGAGTAGTGCCCAAATGC	--	--
	β-actin forward	CATCGCAATGAGCGGTTCC	--	147
	β-actin reverse	ACCGTGTGGCGTAGAGGTC	--	--
Promoter cloning	FATP1 promoter forward	CTACTGTGGTGGGCACTTG	-1856/-1837	2046
	FATP1 promoter reverse	TTGTTCCCTGGCTGACCTGGAG	+168/+190	--
	FATP1 promoter forward 1	GAGCTC TACTGTGGTGGGCACTTG	-1856/-1837	2046
	FATP1 promoter forward 2	GAGCTC CAATGTGCAGAGGTGAGAG	-1558/-1540	1748
	FATP1 promoter forward 3	GAGCTC CGCTAGTGTTAAAGAACCTG	-1261/-1241	1451
	FATP1 promoter forward 4	GAGCTC TCCACTGGAGGAGTACTG	-955/-938	1145
	FATP1 promoter forward 5	GAGCTC AAGGCCACAGGATGGGAGGAGAAAG	-640/-616	830
	FATP1 promoter forward 6	GAGCTC CGCAGAGTGCAAGCCTCAG	-387/-368	577
	FATP1 promoter forward 7	GAGCTC AGAGCTGAGAAGGTCGGCCAAG	-96/-75	286
	FATP1 promoter reverse 1	CTCGAG TGTTCCCTGGCTGACCTGGAG	+168/+190	--
5'RACE	FATP1-GSP1	GAAGTGGTCCGAGAACGGGTTGAGG	--	948
	FATP1-GSP2	CGCATAGTCGTGCTTTCCGGCTTC	--	531
Site-directed mutagenesis	mKLF15 forward	CCAAGCAGGAAAGAAACAAGCA CC GTGGGATAGGCAGGGGGG	-79/-36	286
	mKLF15 reverse	CCCCCTGCCTATCCCAC GG GTGCTTGTTCCTTCTGCTTGG	--	--
	mPPARγ forward	GATAGGCAGGGGGC ATTA GTAGGGGAGCTTGAG	-50/-15	286
	mPPARγ reverse	CTCCAAGCTCCCC TA ATGCCCTGCCTATC	--	--
EMSA	KLF15 forward	CCAAGCAGGAAAGAAACAAGCA GGG GTGGGATAGGCAGGGGGG	-79/-36	286
	KLF15 reverse	CCCCCTGCCTATCCA CC CTGCTTGTTCCTTCTGCTTGG	--	--